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Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: Mon Sep 10 14:22:55 EDT 2007

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Reviewer Comments:

<210> 3

<211> 297

<212> DNA

<213> Synthetic construct

<220>

<223> CXCL8-1B3 coding sequence

<400> 3

The above <213> response for sequence id# 3 is invalid, please correct the remaining sequences. FYI, the above <213> response can be inserted into section <220> - <223> as a response.

****

Validated By CRFValidator v 1.0.3

Application No: 10573726 Version No: 1.0

Input Set:

Output Set:

Started: 2007-08-28 12:12:14.845
Finished: 2007-08-28 12:12:15.124
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 279 ms
Total Warnings: 4
Total Errors: 0
No. of SeqIDs Defined: 6
Actual SeqID Count: 6

Error code	Error Description
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W 402	Undefined organism found in <213> in SEQ ID (4)
W 402	Undefined organism found in <213> in SEQ ID (5)
W 402	Undefined organism found in <213> in SEQ ID (6)

SEQUENCE LISTING

<110> Applied Research Systems ARS Holding N.V.

<120> NOVEL CXCL8 ANTAGONISTS

<130> WO932

<140> 10573726
<141> 2007-08-28

<160> 6

<170> PatentIn version 3.1

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<212> DNA
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<223> Human CXCL8 coding sequence

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tccaaacctt tccaccccaa atttatcaaa gaactgagag tgattgagag tggaccacac 180

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<213> Homo sapiens

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<223> Mature human CXCL8

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20 25 30

His Cys Ala Asn Thr Glu Ile Ile Val Lys Leu Ser Asp Gly Arg Glu
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Leu Cys Leu Asp Pro Lys Glu Asn Trp Val Gln Arg Val Val Glu Lys
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Phe Leu Lys Arg Ala Glu Asn Ser
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tccaaacctt tccaccccaa atttatcaaa gaactgagag tgattgagag tggaccacac 180

tgcgccaaca cagaaattat tgtaaagctt tctgatggaa gagagctctg tctggacccc 240

aaggaaaact gggtgcaggc ggttgtggag gcgtttttgg cgagggctga gaattca 297

<210> 4

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<212> PRT

<213> Synthetic construct

<220>

<223> Mature CXCL8-1B3

<400> 4

Ser Ala Lys Glu Leu Arg Cys Gln Cys Ile Lys Thr Tyr Ser Lys Pro
1 5 10 15

Phe His Pro Lys Phe Ile Lys Glu Leu Arg Val Ile Glu Ser Gly Pro
20 25 30

His Cys Ala Asn Thr Glu Ile Ile Val Lys Leu Ser Asp Gly Arg Glu
35 40 45

Leu Cys Leu Asp Pro Lys Glu Asn Trp Val Gln Ala Val Val Glu Ala
50 55 60

Phe Leu Ala Arg Ala Glu Asn Ser
65 70

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<212> DNA

<213> Synthetic construct

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tgcgccaaca cagaaattat tgtaaagctt tctgatggaa gagagctctg tctggacccc 240

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<212> PRT

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<400> 6

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Ser Ala Lys Glu Leu Arg Cys Gln Cys Ile Lys Thr Tyr Ser Lys Pro
1               5                   10                  15

Phe His Pro Lys Phe Ile Lys Glu Leu Arg Val Ile Glu Ser Gly Pro
            20                  25                  30

His Cys Ala Asn Thr Glu Ile Ile Val Lys Leu Ser Asp Gly Arg Glu
        35                  40                  45

Leu Cys Leu Asp Pro Lys Glu Asn Trp Val Gln Arg Val Val Glu Ala
    50                  55                  60

Phe Leu Ala Ala Ala Glu Asn Ser
65                  70
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